

GenCore version 5.1.6
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OM nucleic - nucleic search, using sw model

Run on: May 4, 2005, 12:02:42 ; Search time 254.353

Seconds

(without alignments)

1814.018 Million cell

updates/sec

Title: US-09-300-482-27

Perfect score: 229

Sequence: 1

cagaccttattttttctgtc.....cttcaatttatggggaacaa 229

Scoring table: IDENTITY_NUC

Gapop 10.0 , Gapext 1.0

Searched: 9369276 seqs, 1007425964 residues

Total number of hits satisfying chosen parameters: 18738552

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Pending_Patents_NA_New:*

- 1: /cgn2_6/ptodata/1/pna/PCT_NEW_COMB.seq:*
- 2: /cgn2_6/ptodata/1/pna/PCT_NEW_COMB.seq2:*
- 3: /cgn2_6/ptodata/1/pna/US06_NEW_COMB.seq:*
- 4: /cgn2_6/ptodata/1/pna/US07_NEW_COMB.seq:*
- 5: /cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq:*
- 6: /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*
- 7: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq:*
- 8: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq2:*
- 9: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq3:*
- 10: /cgn2_6/ptodata/1/pna/US11_NEW_COMB.seq:*
- 11: /cgn2_6/ptodata/1/pna/US11_NEW_COMB.seq2:*
- 12: /cgn2_6/ptodata/1/pna/US11_NEW_COMB.seq3:*
- 13: /cgn2_6/ptodata/1/pna/US60_NEW_COMB.seq:*